RT² Profiler PCR Array (Rotor-Gene® Format) Mouse Epigenetic Chromatin Remodeling Factors

Cat. no. 330231 PAMM-086ZR

For pathway expression analysis

| Format | For use with the following real-time cyclers | | |
|-------------------------------------|----------------------------------------------|--|--|
| RT ² Profiler PCR Array, | Rotor-Gene Q, other Rotor-Gene cyclers | | |
| Format R | | | |

Description

The Mouse Epigenetic Chromatin Remodeling Factors RT² Profiler PCR Array profiles the expression of 84 key genes involved in recognizing chromatin modifications and remodeling chromatin to regulate chromatin accessibility and therefore gene expression. Many of these genes contain the chromodomain responsible for ATP hydrolysis-dependent chromatin reorganization. Chromodomain proteins either activate or repress transcription depending on their additional domains and the proteins they recruit. Chromobox domain proteins recognize methylated histone lysine residues and mediate transcriptional repression. Bromodomains and plant homeodomains (PHD) respectively bind acetylated and methylated histone lysine residues and seem to work cooperatively. Other genes on this array, such as the Inhibitor of Growth (ING) family, associate with and modulate histone acetyltransferase and deacetylase complex activity, while still others directly bind methylated CpG DNA. During the development of stem cells to terminally differentiated cells, altered expression occurs for many of these genes to control chromatin dynamics. These genes also exhibit different expression profiles in tumor cells relative to normal cells, suggesting a role for chromatin modification and remodeling in oncogenesis. Using real-time PCR, you can easily and reliably analyze the expression of a focused panel of genes involved in epigenetic chromatin remodeling with this array.

For further details, consult the RT² Profiler PCR Array Handbook.

Shipping and storage

RT² Profiler PCR Arrays in the Rotor-Gene format are shipped at ambient temperature, on dry ice, or blue ice packs depending on destination and accompanying products.

For long term storage, keep plates at -20°C.

Note: Ensure that you have the correct RT² Profiler PCR Array format for your real-time cycler (see table above).

Note: Open the package and store the products appropriately immediately on receipt.



Array layout

The 96 real-time assays in the Rotor-Gene format are located in wells 1–96 of the Rotor-Disc[™] (plate A1–A12=Rotor-Disc 1–12, plate B1–B12=Rotor-Disc 13–24, etc.). To maintain data analysis compatibility, wells 97–100 do not contain real-time assays but will contain master mix to account for weight balance.

Gene table: RT² Profiler PCR Array

| Position | UniGene | GenBank | Symbol | Description | |
|------------|-----------------------|--------------|--------|------------------------------------------------|--|
| A01 | Mm.22478 | NM 001080819 | Arid1a | AT rich interactive domain 1A (SWI-like) | |
| A02 | Mm.17166 | NM 175251 | Arid2 | AT rich interactive domain 2 (ARID, RFX-like) | |
| A03 | Mm.330677 | NM 001039939 | Asxl1 | Additional sex combs like 1 (Drosophila) | |
| A04 | Mm.485468 | NM 013815 | Baz1a | Bromodomain adjacent to zinc finger domain 1A | |
| A05 | Mm.40331 | NM 011714 | Baz1b | Bromodomain adjacent to zinc finger domain, 1B | |
| A06 | Mm.252213 | NM 054078 | Baz2a | Bromodomain adjacent to zinc finger domain, 2A | |
| A07 | Mm.228071 | NM 001001182 | Baz2b | Bromodomain adjacent to zinc finger domain, 2B | |
| A08 | Mm.289584 | NM 007552 | Bmi1 | Bmi1 polycomb ring finger oncogene | |
| A09 | Mm.343986 | NM 176850 | Bptf | Bromodomain PHD finger transcription factor | |
| A10 | Mm.254438 | NM 001033274 | Brd1 | Bromodomain containing 1 | |
| A11 | Mm.3444 | NM 010238 | Brd2 | Bromodomain containing 2 | |
| A12 | Mm.28721 | NM 023336 | Brd3 | Bromodomain containing 3 | |
| B01 | Mm.253518 | NM 198094 | Brd4 | Bromodomain containing 4 | |
| B02 | Mm.5400 | NM 012047 | Brd7 | Bromodomain containing 7 | |
| B03 | Mm.411740 | NM 030147 | Brd8 | Bromodomain containing 8 | |
| B04 | Mm.182836 | NM 054054 | Brdt | Bromodomain, testis-specific | |
| B05 | Mm.356059 | NM 030178 | Brpf1 | Bromodomain and PHD finger containing, 1 | |
| B05 | Mm.291877 | NM 001081315 | Brpf3 | Bromodomain and PHD finger containing, 7 | |
| B07 | Mm.240871 | NM 145125 | Brwd1 | Bromodomain and WD repeat domain containing 1 | |
| B07 | Mm.100112 | NM 001081477 | Brwd3 | | |
| B08 B09 | Mm.100112 Mm.29055 | NM 007622 | Cbx1 | Bromodomain and WD repeat domain containing 3 | |
| B10 | Mm.29055 Mm.14547 | | Cbx1 | Chromobox homolog 1 (Drosophila HP1 beta) | |
| | | NM_007623 | | Chromobox homolog 2 (Drosophila Pc class) | |
| B11 | Mm.280968 | NM_007624 | Cbx3 | Chromobox homolog 3 (Drosophila HP1 gamma) | |
| B12 | Mm.268070 | NM_007625 | Cbx4 | Chromobox homolog 4 (Drosophila Pc class) | |
| C01 | Mm.262059 | NM_007626 | Cbx5 | Chromobox homolog 5 (Drosophila HP1a) | |
| C02 | Mm.38438 | NM_028763 | Cbx6 | Chromobox homolog 6 | |
| C03 | Mm.323442 | NM_144811 | Cbx7 | Chromobox homolog 7 | |
| C04 | Mm.99953 | NM_013926 | Cbx8 | Chromobox homolog 8 (Drosophila Pc class) | |
| C05 | Mm.29002 | NM_009881 | Cdyl | Chromodomain protein, Y chromosome-like | |
| C06 | Mm.482911 | NM_029441 | Cdyl2 | Chromodomain protein, Y chromosome-like 2 | |
| C07 | Mm.8137 | NM_007690 | Chd1 | Chromodomain helicase DNA binding protein 1 | |
| C08 | Mm.34955 | NM_001081345 | Chd2 | Chromodomain helicase DNA binding protein 2 | |
| C09 | Mm.178246 | NM_146019 | Chd3 | Chromodomain helicase DNA binding protein 3 | |
| C10 | Mm.333388 | NM_145979 | Chd4 | Chromodomain helicase DNA binding protein 4 | |
| C11 | Mm.40192 | NM_001081376 | Chd5 | Chromodomain helicase DNA binding protein 5 | |
| C12 | Mm.122738 | NM_173368 | Chd6 | Chromodomain helicase DNA binding protein 6 | |
| D01 | Mm.138792 | NM_001081417 | Chd7 | Chromodomain helicase DNA binding protein 7 | |
| D02 | Mm.289934 | NM_201637 | Chd8 | Chromodomain helicase DNA binding protein 8 | |
| D03 | Mm.100615 | NM_177224 | Chd9 | Chromodomain helicase DNA binding protein 9 | |
| D04 | Mm.7286 | NM_013502 | Ctbp1 | C-terminal binding protein 1 | |
| D05 | Mm.246240 | NM_009980 | Ctbp2 | C-terminal binding protein 2 | |
| D06 | Mm.269474 | NM_181322 | Ctcf | CCCTC-binding factor | |
| D07 | Mm.380914 | NM_021876 | Eed | Embryonic ectoderm development | |
| D08 | Mm.246688 | NM_007971 | Ezh2 | Enhancer of zeste homolog 2 (Drosophila) | |
| D09 | Mm.24570 | NM_172162 | Hinfp | Histone H4 transcription factor | |
| D10 | Mm.25709 | NM_011919 | Ing1 | Inhibitor of growth family, member 1 | |
| D11 | Mm.430733 | NM 023503 | Ing2 | Inhibitor of growth family, member 2 | |
| D12 | Mm.39999 | NM_023626 | Ing3 | Inhibitor of growth family, member 3 | |
| E01 | Mm.262547 | NM 133345 | Ing4 | Inhibitor of growth family, member 4 | |
| E02 | Mm.64065 | NM 025454 | Ing5 | Inhibitor of growth family, member 5 | |
| E03 | Mm.330496 | NM 026574 | Ino80 | INO80 homolog (S. cerevisiae) | |
| E04 | Mm.22522 | NM 013594 | Mbd1 | Methyl-CpG binding domain protein 1 | |
| E05 | Mm.7142 | NM 013595 | Mbd3 | Methyl-CpG binding domain protein 3 | |
| E06 | Mm.259308 | NM 010774 | Mbd4 | Methyl-CpG binding domain protein 4 | |
| E07 | Mm.131408 | NM 010788 | Mecp2 | Methyl CpG binding protein 2 | |
| E08 | Mm.212577 | NM 054081 | Mta 1 | Metastasis associated 1 | |
| E09 | Mm.25339 | NM 011842 | Mta2 | Metastasis-associated gene family, member 2 | |
| LU7 | 14111.23337 | 14/4/_011042 | MIUZ | melusiusis-ussociuleu gene lunniy, member 2 | |

| Position | UniGene | GenBank | Symbol | Description | |
|----------|----------------------------|--------------|----------|-----------------------------------------------------------------------------|--|
| E10 | Mm.336898 | NM_008668 | Nab2 | Ngfi-A binding protein 2 | |
| E11 | Mm.12964 | NM_008739 | Nsd1 | Nuclear receptor-binding SET-domain protein 1 | |
| E12 | Mm.12261 | NM_197992 | Pcgf1 | Polycomb group ring finger 1 | |
| F01 | Mm.2418 | NM_009545 | Pcgf2 | Polycomb group ring finger 2 | |
| F02 | Mm.261218 | NM_172716 | Pcgf3 | Polycomb group ring finger 3 | |
| F03 | Mm.259180 | NM_029508 | Pcgf5 | Polycomb group ring finger 5 | |
| F04 | Mm.35413 | NM_027654 | Pcgf6 | Polycomb group ring finger 6 | |
| F05 | Mm.6822 | NM_007905 | Phc1 | Polyhomeotic-like 1 (Drosophila) | |
| F06 | Mm.259103 | NM_018774 | Phc2 | Polyhomeotic-like 2 (Drosophila) | |
| F07 | Mm.480590 | NM_009343 | Phf1 | PHD finger protein 1 | |
| F08 | Mm.25582 | NM_172705 | Phf13 | PHD finger protein 13 | |
| F09 | Mm.212554 | NM_011078 | Phf2 | PHD finger protein 2 | |
| F10 | Mm.101022 | NM_001081166 | Phf21b | PHD finger protein 21B | |
| F11 | Mm.194486 | NM_001081080 | Phf3 | PHD finger protein 3 | |
| F12 | Mm.271715 | NM_026737 | Phf5a | PHD finger protein 5A | |
| G01 | Mm.26870 | NM_027642 | Phf6 | PHD finger protein 6 | |
| G02 | Mm.5348 | NM_027949 | Phf7 | PHD finger protein 7 | |
| G03 | Mm.20343 | NM_009066 | Ring1 | Ring finger protein 1 | |
| G04 | Mm.482250 | NM_011277 | Rnf2 | Ring finger protein 2 | |
| G05 | Mm.313303 | NM_011416 | Smarca2 | SWI/SNF related, matrix associated, actin dependent regulator of chromatin, | |
| G05 | | | | subfamily a, member 2 | |
| G06 | Mm.286593 | NM_011417 | Smarca4 | SWI/SNF related, matrix associated, actin dependent regulator of chromatin, | |
| G00 | | | | subfamily a, member 4 | |
| C07 | G07 Mm.246803 NM_053124 Sr | | Smarca5 | SWI/SNF related, matrix associated, actin dependent regulator of chromatin, | |
| G07 | | | Smarcas | subfamily a, member 5 | |
| G08 | Mm.299906 | NM_019763 | Spen | SPEN homolog, transcriptional regulator (Drosophila) | |
| G09 | Mm.283410 | NM_199196 | Suz12 | Suppressor of zeste 12 homolog (Drosophila) | |
| G10 | Mm.314056 | NM_009054 | Trim27 | Tripartite motif-containing 27 | |
| G11 | Mm.229323 | NM_172255 | Wdr11 | WD repeat domain 11 | |
| G12 | Mm.227598 | NM_027230 | Zmynd8 | Zinc finger, MYND-type containing 8 | |
| H01 | Mm.328431 | NM_007393 | Actb | Actin, beta | |
| H02 | Mm.163 | NM_009735 | B2m | Beta-2 microglobulin | |
| H03 | Mm.343110 | NM_008084 | Gapdh | Glyceraldehyde-3-phosphate dehydrogenase | |
| H04 | Mm.3317 | NM_010368 | Gusb | Glucuronidase, beta | |
| H05 | Mm.2180 | NM_008302 | Hsp90ab1 | Heat shock protein 90 alpha (cytosolic), class B member 1 | |
| H06 | N/A | SA_00106 | MGDC | Mouse Genomic DNA Contamination | |
| H07 | N/A | SA_00104 | RTC | Reverse Transcription Control | |
| H08 | N/A | SA_00104 | RTC | Reverse Transcription Control | |
| H09 | N/A | SA_00104 | RTC | Reverse Transcription Control | |
| H10 | N/A | SA_00103 | PPC | Positive PCR Control | |
| H11 | N/A | SA_00103 | PPC | Positive PCR Control | |
| H12 | N/A | SA_00103 | PPC | Positive PCR Control | |

Related products

For optimal performance, RT² Profiler PCR Arrays should be used together with the RT² First Strand Kit for cDNA synthesis and RT² SYBR[®] Green qPCR Mastermixes for PCR.

| Product | Contents | Cat. no. |
|--------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------|----------|
| RT ² First Strand Kit (12) | Enzymes and reagents for cDNA synthesis | 330401 |
| RT ² SYBR Green ROX [™] FAST Mastermix (2)* | For 2 x 96 assays in 96-well plates; suitable for use with the Rotor-Gene Q and other Rotor-Gene cyclers | 330620 |

^{*} Larger kit sizes available; please inquire.

RT² Profiler PCR Array products are intended for molecular biology applications. These products are not intended for the diagnosis, prevention, or treatment of a disease.

For up-to-date licensing information and product-specific disclaimers, see the respective QIAGEN kit handbook or user manual. QIAGEN kit handbooks and user manuals are available at www.qiagen. com or can be requested from QIAGEN Technical Services or your local distributor.

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